



Supplementary Materials

Table S1. Number of samples collected and of pigs and farm workers colonized with *S. aureus* in each farm investigated in the study.

| Farm | Region | Collection date | Pig (n) | | Farm work (n) | |
|----------------|--------|-----------------|--------------|-----------|---------------|-----------|
| | | | Investigated | Colonized | Investigated | Colonized |
| A | II | 01/2014 | 10 | 2 | 2 | 0 |
| B | II | 01/2014 | 10 | 2 | 2 | 0 |
| C | I | 01/2014 | 10 | 0 | 1 | 0 |
| D | II | 01/2014 | 10 | 0 | 1 | 0 |
| E | II | 01/2014 | 10 | 2 | - | - |
| F | III | 01/2014 | 12 | 0 | - | - |
| G ¹ | I | 01/2014 | 7 | 1 | - | - |
| H | IV | 01/2014 | 10 | 0 | - | - |
| I | IV | 01/2014 | 16 | 0 | - | - |
| J | IV | 01/2014 | 5 | 0 | - | - |
| K | I | 12/2015 | 21 | 1 | 5 | 2 |
| L | III | 11/2016 | 20 | 4 | 4 | 1 |
| M | I | 05/2019 | 10 | 1 | 1 | 0 |
| N | III | 06/2019 | 25 | 1 | 5 | 3 |
| O ¹ | I | 08/2019 | 23 | 0 | 6 | 1 |
| P | I | 11/2019 | 31 | 0 | 4 | 0 |
| Total | | | 230 | 14 | 27* | 7 |

¹Farms A and B and farms G and O are localized in the same cities. * Four farms (A, B, C and D) were attended by the same veterinarian, and his sample was collected only once. Samples from animals and farm workers were collected on the same date on these four farms.

Table S2. Antimicrobial resistance patterns of multidrug-resistant *Staphylococcus aureus* strains from pigs and farm workers in pig farms from Rio de Janeiro state.

| Region | Farm | Source | Host | Antimicrobial resistance pattern ² | |
|--------|------|--------|--------------------|---|---|
| I | K | Pig | SN105 | Chl (I), Cip, Cli, Ery, Nor, Pen, Sut (I), Tet | |
| | | Human | HSN10 | iCli, Ery, Pen, Tet | |
| | O | Human | HSN21 | iCli, Ery, Gen, Pen | |
| II | B | Pig | SN15 | Chl, Cip, Cli, Ery, Pen, Tet | |
| | E | Pig | SN51 | Cef, Cip, Cli, Ery, Nor, Pen, Sut, Tet | |
| | | Pig | SN52 | Cef, Cip, Cli, Ery, Nor, Pen, Tet ³ | |
| III | L | Pig | SN128 | Chl (I), Cip, Cli, Ery, Gen, Nor, Pen, Tet | |
| | | Pig | SN143 | Chl (I), Cip, Cli, Ery, Gen, Nor, Pen, Sut, Tet | |
| | | Pig | SN144 | Chl (I), Cip (I), Cli, Ery, Pen, Tet | |
| | | Pig | SN145 ¹ | Chl, Cip, Cli, Ery, Gen, Nor, Pen, Tet Chl, Cli, Ery, Pen, Sut, Tet | |
| | N | Human | HSN12 ¹ | Chl, Cip, Cli, Ery, Gen, Nor, Pen, Tet Chl, Cip, Cli, Ery, Nor, Pen, Tet Chl (I), Cip, Cli, Ery, Nor, Pen, Tet | |
| | | | Pig | SN182 ¹ | Chl, Cip, Cli, Tet Chl, Cip, Cli, Ery, Pen, Tet⁴ |
| | | | Human | HSN16 | Cip, Cli, Pen, Tet |
| | | Human | HSN18 | Chl, Cip, Cli, Ery, Pen, Tet⁴ | |

¹Antimicrobial resistance pattern of more than one bacterial strain per pig or human; ²Cef: ceftiofur (detection of MRSA, methicillin-resistant *Staphylococcus aureus*), Cip: ciprofloxacin, Cip (I): intermediate to ciprofloxacin, Chl: chloramphenicol; Chl (I): intermediate to chloramphenicol; Cli: clindamycin, iCli: inducible clindamycin resistance, Ery: erythromycin, Gen: gentamicin, Nor: norfloxacin, Pen: penicillin G, Sut: sulfamethoxazole-trimethoprim, Tet: tetracycline; ³MRSA; ⁴OS-MRSA (oxacillin-susceptible MRSA). Indistinguishable antimicrobial patterns of pig and human from the same farm are in bold.